MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	<b>APONTS</b> HGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of Xenopus cerberus protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the Xenopus organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTCCCAC	CAAGTCGCTC	. PCSSSCSCWC	010000000			
CTTAAGGGT	GTTCAGCGAG	. WOUNTACHCIG	CAGGGTCTAG	ATATCATACA	atgttactaa	60
	GTTCAGCGAG	ICITIGICAC	GTCCCAGATC	TATAGTATGT	TACAATGATT	
ATGTACTCAC	GATCTGTATT	<b>ልጥ</b> ርርጥርጥርርር	##C#C\$ \$ #C\$			
TACATGAGTO	CTAGACATAA	TACCACACAC	11G1GAATGA	TGGAGCAGGA	AAACACTCAG	120
	•	* NOCHONCOG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
AAGGACGAGA	AAGGACAAAA	ACATATTCAC	TT3 3 C 3 C C 3 C	10000000000		
TTCCTGCTCT	TTCCTGTTTT	TGTATAACTC	A A TOTO COCOCO	AGGTTACTTC	AGAAAAGAAA	180
		-01111111111111111111111111111111111111	ANTIGICGIC	TCCAATGAAG	TCTTTTCTTT	
GAGGAGCACG	TAGGAGCAAG	ATTCTGCTGG	<b>でになるでものである</b>	ACCTCTTCA	633,66663	
CTCCTCGTGC	ATCCTCGTTC	TAAGACGACC	ACTTATCATE	TCCACAACTA	GAACCCCACA	240
				ICCAGAACIA	CTTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AACTATTTCA	TTCCACCACA	ACACAMACA A	200
AACCCGTACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	A A COTTOTO	MCACATACAA	300
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATCCAAACA	200
TGTCTTTTCT	CGGTCTGTAC	TTGTTTCAGT	TCGAAAAGAG	TTGTCAACGG	CAT GOVERNO	360
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	<b>ፕ</b> ልጥጥጥጥርርጥ	<b>CCCCCののののの</b> の	420
TTTCACGTTC	TTCTTTTCGA	ATGTTACCAA	GATCTTCCTT	ATAAAAAGGA	GCGCCAACAA	420
TTGATAAAAG	AAATACAGAG	GTTACTGAAA	AGCCTGGTGC	CAAGATGTTC	TGGDDCDDTT	480
AACTATTTTC	TTTATGTCTC	CAATGACTTT	TCGGACCACG	GTTCTACAAG	ACCTTGTTAA	400
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCGGT	ACCGTCATTT	CGTGTCCTTT	340
TAATGAAAGA	AGCTTGCAAA	ACCTTGTTTT	TCACTCAGAA	TATTGTACAT	GAAAACTGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	•••
ACAGGATGGT	GATACAGAAC	AATCTGTGCT	TTGGTAAATG	CATCTCTCTC	CATGTTCCAA	660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
ATCAGCAAGA	TCGACGAAAT	ACTTGTTCCC	ATTGCTTGCC	GTCCAAATTT	ACCCTGAACC	720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
ACCECT COOR	<b>6110000</b>					
TECNETOCO	GAATTGTACT	GGATCTAAGA	ATGTAGTAAA	GGTTGTCATG	ATGGTAGAGG	780
1GGAC1GCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATCCACCTC	5C11C05015	***				
TTACCTCCAC	TGAAGCTCAT	AAGAGCAACT	TCCACCAAAC	TGCACAGTTT	AACATGGATA	840
I INCOIGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
CATCTACTAC	CCTCCACCAM	<b>5111001000</b>				
GTAGATGATG	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TGGAAATGCC	Cttttgttgg	900
	GGACGTGGTA	ATTICCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
AATATTTGTT	ACATACTATO	<b>ころかいかきょうこ</b> つ	1 mm 1 mc			
TTATAAACAA	ACATACTATG TGTATGATAC	CTACAMMMCC	ATTATGTTGC	CTTCTATTTC	ATATAACCAC	960
	TGTATGATAC	O-WOUT I I CO	TARTACARCG	GAAGATAAAG	TATATTGGTG	
ATGGAATAAG	GATTGTATGA	<b>ልጥተልጥል</b> ልጥጥጻ	8C8880CCC			. —
TACCTTATTC	CTAACATACT	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	MCMMATGGCA	TTTTGTGTAA	CATGCAAGAT	1020
		INTINAL	IGITTACCGT	AAAACACATT	GTACGTTCTA	

CTCTGTTCCA	TCAGTTGCAA	GATAAAAGGC	AATATTTGTT	TGACTTTTTT	TCTACAAAAT	1080
GAGACAAGGT	AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA	ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
CTTATGGGTT	TATATACTAT	TCTATTACCC	CAGTTTTGAC	AATTCCCCAT	TACATTATTA	
AGGGACTAAG	TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTC	AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
TGGTCACCTG	TTTAAAAGCA	AACATCTTAT	TGGTTGCTAT	GGGTTACTGC	TTCTGGGCAA	1260
ACCAGTGGAC	AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC	TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTCAT	1320
TTACACACGG	AGTATCCCCC	CAATCACACA	ACACATGACT	TATTTAACAT	AAATAAAGTA	
TGTTACAAAA ACAATGTTTT						

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSRTRKVDSL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	NHLHHSTQAN	60
AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSVCER	ARAGCEPILI	120
KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
KPMKATQKTY	LKNNYNYVIR	AKVKEVKVKC	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
SGCLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKKVKRWD	QKLRRPRKSK	300
DPVAPIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

C	CACAGGA CTCCTGGCAG	NCCDCN NDCC	mm> CCCCm> m	CC 3 BBBCCBB	60
	GTGTCCT GAGGACCGTC				60
011111000000000000000000000000000000000	diodiccoic	ICCACTIACC	MICOGGNIA	CCIAMCOM	
TGTTGATTTT GAC	ACATGAT TGATTGCTTT	CAGATAGGAT	TGAAGGACTT	GGATTTTTAT	120
ACAACTAAAA CTG	STGTACTA ACTAACGAAA	GTCTATCCTA	ACTTCCTGAA	CCTAAAAATA	
	TTTAAAT TATCTGAGTA				180
GATTAAGACG TGA	AAATTTA ATAGACTCAT	TAACAAGTAA	AACATAACCT	ACCCTGATTT	
GATAAACTTA ACT	CCTTGCT TTTGACTTGC	CCATAAACTA	TAAGGTGGGG	TGAGTTGTAG	240
	GGAACGA AAACTGAACG	*			
TTGCTTTTAC ATG	STGCCCAG ATTTTCCCTG	TATTCCCTGT	ATTCCCTCTA	AAGTAAGCCT	300
AACGAAAATG TAC	CACGGGTC TAAAAGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCGGA	
1010151010 055					2.50
	GGGCAGA ATAACAATGT				360
TGTGTATGTC CAA	ACCCGTCT TATTGTTACA	GAGCTTGTTC	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT ACC	TGGACTG GCGCTTCTCT	TATTACCCAA	TGCTTACTGT	GCTTCGTGTG	420
	SACCTGAC CGCGAAGAGA				
AGCCTGTGCG GAT	CCCCATG TGCAAATCTA	TGCCATGGAA	CATGACCAAG	ATGCCCAACC	480
TCGGACACGC CTA	AGGGGTAC ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
	SCACTCAA GCCAATGCCA				540
TAGAGGTGGT GTC	GTGAGTT CGGTTACGGT	AGGACCGTTA	ACTTGTCAAA	CTTCCAAACG	
TGACCACTGA ATG	STAGCCAG GACCTTTTG1	TCTTTCTGTG	TGCCATGTAT	GCCCCCATTT	600
	ATCGGTC CTGGAAAACA				-
GTACCATCGA TTT	CCAGCAT GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGGCCA	660
CATGGTAGCT AAR	AGGTCGTA CTTGGTTAAT	TCGGAACGTT	CAGGCACACG	CTTTCCCGGT	
	AGCCCATT CTCATAAAG1				720
CCCGGCCGAC AC	icgggtaa gagtatttci	1 16GCCGTGTG	AACCGGTCTC	TCGGACCGTA	
GTGAAGAGCT GCC	CCGTATAT GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
	GCATATA CTGTCTCCTC				
TGGAACAAGG AAG	CAGATTCA ATGCCAGACT	TCTCCATGGA	TTCAAACAAT	GGAAATTGCG	840
ACCTTGTTCC TTC	STCTAAGT TACGGTCTG	AGAGGTACCT	AAGTTTGTTA	CCTTTAACGC	
	AGCACTGT AAATGCAAG				900
CTTCGCCGTC CC	ICGTGACA TTTACGTTC	GGTACTTCCG	TTGGGTTTTC	TGCATAGAGT	
AGAATAATTA CAI	ATTATGTA ATCAGAGCAI	AAGTGAAAGA	GGTGAAAGTG	AAATGCCACG	960
	PAATACAT TAGTCTCGT				200
•					
ACGCAACAGC AA	TTGTGGAA GTAAAGGAG	A TTCTCAAGTC	TTCCCTAGTG	AACATTCCTA	1020
TGCGTTGTCG TT	AACACCTT CATTTCCTC	r aagagttcag	AAGGGATCAC	TTGTAAGGAT	

,	AGACACAGT	GACACTGTAC	ACCAACTCAG	GCTGCTTGTG	CCCCAGCTT	GTTGCCAATG	1080
				CGACGAACAC			
				AAGAGCGTAC			1140
1	CCTTATGTA	TTAATACCCG	ATACTTCTGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
_						-000180111	1000
				TTGCTAAGAA			1200
(	TAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
7	このでもないの	TCCCAGGAAA	ACCAAACACC	<b>CCCTCCCTCC</b>	AATTCCCAAC	AAAAACAGCA	1260
				GGCACCGAGG			
•	CGARGC1GC	AGGGICCIII	1031110193	GOUNCOGNOG	11111000110		
,	TTCCAGACA	AGCGCGTAGT	TAGACTAACG	GAAAGGTGTA	TGGAAACTCT	ATGGACTTTG	1320
				CTTTCCACAT			
•							
1	AAACTAAGAT	TTGCATTGTT	GGAAGAGCAA	AAAAGAAATT	GCACTACAGC	ACGTTATATT	1380
•	<b>TTGATTCTA</b>	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
						TTCTTTTTTT	1440
(	GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAA	
							1500
							1500
1	<b>AATATTGATA</b>	TAAACGTGCA	CAAGGGTCCG	TTAACAAAAT	AAGTTGAAGG	TCACTGTCTC	
		> momomo> 00	0777707700	#C33##C3##	mcmcsmcssc	TAATGGTGAC	1560
						ATTACCACTG	1300
•	GTCACTGACT	TACAGAGTCG	GATTICITCS	AGIIAAGIAA	AGACIAGIIG	ATTACCACTO	
	<b>あるこれの中でする</b> あ	<b>ተል</b> ርተተርርርርል	A A CTCA A CTA	<b>አ</b> ተተርርል አተርር	TANATCAGAG	AAAAGTTGAC	1620
						TTTTCAACTG	
	11 CACABACT	211 GE 100001					
	CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
						ACTAGTGAAA	
						AAATCTAAAT	1740
	<b>GGTAAATT</b> AT	GAAAGTCGTC	AAAATCAATC	TACTGTACAT	CCTACGTGG	TTTAGATTTA	
						AGGTAAATGC	1800
	<b>Taaaata</b> gta	TTTACTTCTC	GACCAAATCI	GACATACCAG	TGACAACCC	TCCATTTACG	
			<b></b>				1860
						AAAAAATAAA 7 TTTTTTTATTT /	T000
	GATGAAACAG	TIMMGACAAA	AIIIITAACG	GALITATITA	INNIICAGG		
	AAAAAAAA	4444					
	TTTTTTTTT						

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL 120 MKQFNNSLIG VRESDGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 180 RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER 240 STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVROLFK INSRTGSVTL 300 EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 360 IPETATKENF IALISTIDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TISTLDRENI AAYSLTVVAE DLGFPSLKTK KYYTVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 480 ARDSDSDQNG KVNYRLVDAK VMGQSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA 540 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD 600 EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKOLNSDHSE DLSIVVAVYD LGRPSLSTNA 660 TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 720 AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENCS VSSNQEQHQQ 780 TGIKHSISVP SYHTSGWHLD NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 840 RRALSSQCRH KPVLNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL

Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the Xenopus organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTCCCAG AGATGAACT	CTTGAGATTG	TTTTAAATGA	CTGCAGGTCT	GGAAGGATTC	60
CTTAAGGGTC TCTACTTGA	GAACTCTAAC	AAAATTTACT	GACGTCCAGA	CCTTCCTAAG	
ACATTGCCAC ACTGTTTCT	GGCATGAAAA	AACTGCAAGT	TTCAACTTTG	TTTTTGGTGC	120
TGTAACGGTG TGACAAAGA					
AACTTTGATT CTTCAAGAT	CTGCTTCTCT	TCAGAGCCAT	TCCAATGCTG	CTGTTGGGAC	180
TTGAAACTAA GAAGTTCTA					
TGATGGTTTT ACAAACAGA	TGTGAAATTG	CCCAGTACTA	CATAGATGAA	GAAGAACCCC	240
ACTACCAAAA TGTTTGTCTC					•
CTGGCACTGT AATTGCAGT	TTGTCACAAC	ACTCCATATT	TAACACTACA	GATATACCTG	300
GACCGTGACA TTAACGTCAG					
CAACCAATTT CCGTCTAATO	AAGCAATTTA	ATAATTCCCT	TATCGGAGTC	CGTGAGAGTG	360
GTTGGTTAAA GGCAGATTA					
ATGGGCAGCT GAGCATCATO					420
TACCCGTCGA CTCGTAGTAC					
ACTGCAACCT GGCTTTGGAS					480
TGACGTTGGA CCGAAACCT					
TGAAAGTGGA GGTGAGAGA	ATTAATGACC	ATAGCCCTCA	CTTTCCCAGT	GAAATAATGC	540
ACTITCACCI CCACTCTCTC					
ATGTGGAGGT GTCTGAAAG	TCCTCTGTGG	GCACCAGGAT	TCCTTTAGAA	ATTGCAATAG	600
TACACCTCCA CAGACTTTC					
ATGAAGATGT TGGGTCCAAG	TCCATCCAGA	ACTTTCAGAT	CTCAAATAAT	AGCCACTTCA	660
TACTTCTACA ACCCAGGTTC					
GCATTGATGT GCTAACCAG	GCAGATGGGG	TGAAATATGC	AGATTTAGTC	TTAATGAGAG	720
CGTAACTACA CGATTGGTC					
AACTGGACAG GGAAATCCAG	CCAACATACA	TAATGGAGCT	ACTAGCAATG	GATGGGGGTG	780
TTGACCTGTC CCTTTAGGTC					
TACCATCACT ATCTGGTACT					840
ATGGTAGTGA TAGACCATG					
GCCCAGTGTT TGAGAGAAG	ACCATTGCTG	TGGACCTAGT	AGAGGATGCT	CCTCTGGGAT	900
CGGGTCACAA ACTCTCTTCC					
ACCTTTGTT GGAGTTACA	GCTACTGACG	ATGATGAAGG	AGTGAATGGA	GAAATTGTTT	960
TGGAAAACAA CCTCAATGT					
ATGGATTCAG CACTTTGGC	TCTCAAGAGG	TACGTCAGCT	ATTTAAAATT	AACTCCAGAA	1020
TACCTAAGTC GTGAAACCG	AGAGTTCTCC	ATGCAGTCGA	TAAATTTTAA	TTGAGGTCTT	

CTGGCAGTGT TACTCTTGAA GACCGTCACA ATGAGAACTT	GGCCAAGTTG CCGGTTCAAC	attttgagac taaaactctg	CAAGCAGACT GTTCGTCTGA	TACGAATTTG ATGCTTAAAC	1080
AGGTACAAGC CCAAGATTTG TCCATGTTCG GGTTCTAAAC	GGCCCCAACC CCGGGGTTGG	CACTGACTGC GTGACTGACG	TACTTGTAAA ATGAACATTT	GTAACTGTTC CATTGACAAG	1140
ATATACTTGA TGTAAATGAT TATATGAACT ACATTTACTA	AATACCCCAG TTATGGGGTC	CCATCACTAT GGTAGTGATA	TACCCCTCTG ATGGGGAGAC	ACTACTGTAA TGATGACATT	1200
ATGCAGGAGT TGCCTATATT TACGTCCTCA ACGGATATAA	CCAGAAACAG GGTCTTTGTC	CCACAAAGGA GGTGTTTCCT	GAACTTTATA CTTGAAATAT	GCTCTGATCA CGAGACTAGT	1260
GCACTACTGA CAGAGCCTCT CGTGATGACT GTCTCGGAGA	GGATCTAATG CCTAGATTAC	GACAAGTTCG CTGTTCAAGC	CTGTACTCTT GACATGAGAA	TATGGACATG ATACCTGTAC	1320
AGCACTTTAA ACTACAGCAA TCGTGAAATT TGATGTCGTT	GCTTATGAGG CGAATACTCC	ACAGTTACAT TGTCAATGTA	GATAGTTACC CTATCAATGG	ACCTCTACTT TGGAGATGAA	1380
TAGACAGGGA AAACATAGCA ATCTGTCCCT TTTGTATCGT					1440
CCTCATTGAA GACCAAAAAG GGAGTAACTT CTGGTTTTTC					1500
CTGTATTTTC TAAACCCCAG GACATAAAAG ATTTGGGGTC					1560
ATATAACTAC AGTGATAGCC TATATTGATG TCACTATCGG					1620
GACTTGTGGA TGCAAAAGTG CTGAACACCT ACGTTTTCAC					1680
ACTCTGGAGT ATTGAGAGCT TGAGACCTCA TAACTCTCGA					1740
TTGAAATTGA AGCTGCAGAC AACTTTAACT TCGACGTCTG					1800
TCAGAATAGT TGATCAAAAT AGTCTTATCA ACTAGTTTTA					1860
GCTCGGGTGA AGTTCTGCTT CGAGCCCACT TCAAGACGAA					1920
AAGCCGAGGA TTCAGATGAA TTCGGCTCCT AAGTCTACTT	GGGCACAACT CCCGTGTTGA	CCCAGCTGTT GGGTCGACAA	CTATACCATA GATATGGTAT	CTGAGAGATC GACTCTCTAG	1980
CAAGCAGATT GTTTGCCATT GTTCGTCTAA CAAACGGTAA					2040
ACTCTGACCA TTCAGAGGAC TGAGACTGGT AAGTCTCCTG					2100
CATTATCCAC CAATGCTACA GTAATAGGTG GTTACGATGT					2160

Fig. 6. (Continuation page 2, SEQ ID NO:6).

AAGTCGTTAT TTTC	GCAACCA TCTGCAGAA	G AGCAGCACCA	GATCGATATG	TCCATTATAT	- 2220
TTCAGCAATA AAA	CGTTGGT AGACGTCTT	C TCGTCGTGGT	CTAGCTATAC	AGGTAATATA	
TCATTGCAGT GCT	GGCTGGT GGTTGTGCT	T TGCTACTTTT	GGCCATCTTT	TTTGTGGCCT	2280
AGTAACGTCA CGAC	CCGACCA CCAACACGA	a acgatgaaaa	CCGGTAGAAA	AAACACCGGA	
001000000000000000000000000000000000000					
GTACTTGTAA AAAG	GAAAGCT GGTGAATTT	A AGCAGGTACC	TGAACAACAC	GGAACATGCA	2340
CATGAACATT TTTC	CTTTCGA CCACTTAAA	T TCGTCCATGG	ACTTGTTGTG	CCTTGTACGT	
1701101100					
ATGAAGAACG CCTG	STTAAGC ACCCCATCT	CCCAGTCGGT	CTCTTCTTCT	TTGTCTCAGT	2400
TACTICITEC GGAC	CAATTCG TGGGGTAGA	G GGGTCAGCCA	GAGAAGAAGA	<b>A</b> ACAGAGTCA	
OBC1080180 0011	000000 1001100				
CACTORCICATE COM	ACTOTOC ATCAATACT	G AATCTGAGAA	TTGCAGCGTG	TCCTCTAACC	2460
GACICAGIAC GGT	rgagagg tagttatga	C TTAGACTCTT	AACGTCGCAC	AGGAGATTGG	
NACACCACCA MONO	*C333C3				
THE TOTAL TOTAL	CAAACA GGCATAAAG	ACTCCATCTC	TGTACCATCT	TATCACACAT	2520
TICICGICGI AGIC	CGTTTGT CCGTATTTC	J TGAGGTAGAG	ACATGGTAGA	ATAGTGTGTA	
<u> </u>	**************************************				
CACCAACCCT CCAC	GACAAT TGTGCAATG	A GCATAAGTGG	ACATTCTCAC	ATGGGGCACA	2580
GACCAACCGI GGAC	CCTGTTA ACACGTTAC	r CGTATTCACC	TGTAAGAGTG	TACCCCGTGT	
サヤカニヤカニカカ ここでき	*C*C#CC				
AATCATCTTT CCAT	ACAGTGG GCAAAGGAG	TAGTGACTTC	AATGACAGTG	ACTCTGATAC	2640
MATCHIGITI CCAI	GTCACC CGTTTCCTC	r ATCACTGAAG	TTACTGTCAC	TGAGACTATG	
TAGTGGAGAA TCAG	GAAAAGA AGAGCATTG		0300030330	00100000	0700
ATCACCTCTT ACTO	CTTTTCT TCTCGTAAC	r CCRCCCMMAC	CROCCRCRRC	CCAGTGCTCA	2700
	JIIIIOI IOIOGIAAC.	CGICGGIIAC	GICCGIGIIC	GGTCACGAGT	
ATACACAGAT GAAT	CAGCAG GGTTCCGAC	TCCCCNTNNC	#3####################################	BCCCB B MCB B	27.60
TATGTGTCTA CTTA	AGTCGTC CCAAGGCTG	T ACCCCATANC	ATARACTOC	MCCCMM1CAA	2760
		. MOGGCIAIIG	ATAMAGICGG	IGGCIIAGII	
CAAGGGTCCA GAAA	ATGGGA ACTGCACAT	CCARTATCAR	AACCCCTATA	<b>CACTCTCTTA</b>	2820
GTTCCCAGGT CTT1	TACCCT TGACGTGTA	CGTTATACTT	TTCCCCATAT	CTCDCDCDDT	2020
				OI GROWINI	
CTCTGTAGCT CCTG	STATATT ACAATACCT	CCATGCAAGA	ATGCCTAACC	TGCACATACC	2880
GAGACATCGA GGAC	CATATAA TGTTATGGA	GGTACGTTCT	TACGGATTGG	ACGTGTATGG	2000
GAACCATACC CTTA	AGAGACC CTTATTACC	A TATCAATAAT	CCTGTTGCTA	ATCGGATGCA	2940
CTTGGTATGG GAAT	CTCTGG GAATAATGG	ATAGTTATTA	GGACAACGAT	TAGCCTACGT	-540
GGCGGAATAT GAAA	GAGATT TAGTCAACA	AAGTGCAACG	TTATCTCCGC	AGAGATCGTC	3000
CCGCCTTATA CTTT	CTCTAA ATCAGTTGT	TTCACGTTGC	AATAGAGGCG	TCTCTAGCAG	4
TAGCAGATAC CAAG	SAATTCA ATTACAGTC	CGCAGATATCA	AGACAGCTTC	ATCCTTCAGA	3060
ATCGTCTATG GTTC	CTTAAGT TAATGTCAG	GCGTCTATAGT	TCTGTCGAAG	TAGGAAGTCT	
AATTGCTACA ACCT	TTTAAT CATTAGGCA	I GCAAGTGAGA	ATGCACAAAG	GCAAGTGCTT	3120
TTAACGATGT TGGA	LAAATTA GTAATCCGT	A CGTTCACTCT	TACGTGTTTC	CGTTCACGAA	
	AATATA TGGAGTCTC				3180
ATCGTACTTT CGAT	TTATAT ACCTCAGAG	G GGAAAGGGAG	ACTACCTACC	CCCCTCTGTG	
	LATATAC AGCTGCTTT				3240
TCCTGTCACG TATT	TATATG TCGACGAAA	G ATAAACGTAA	AGTGAACCCT	TAAAAAACAA	
	PATTTTT CCTGAATTG				3300
AAAAAATGTA TAAA	TAAAAA GGACTTAAC	TACACTGTAA	CAGGACAGTG	GATTGATCGT	

Fig. 6. (Continuation page 3, SEQ ID NO:6).

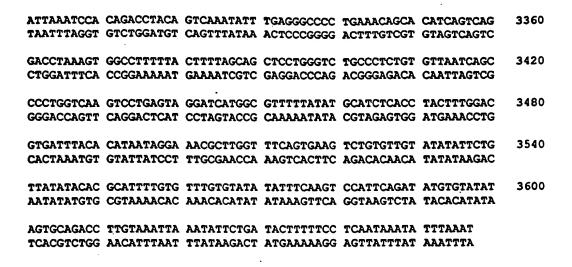
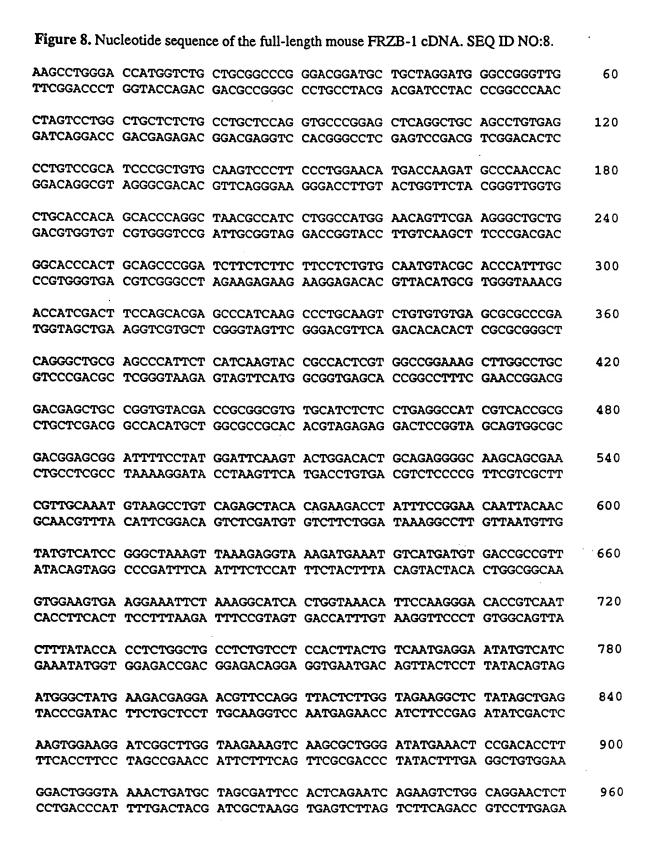


Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180
KPVRATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300
TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.



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	CAGCACGCAG GTCGTGCGTC					1020
	GGCGCTGGTG					1080
	CCGCGACCAC					
	CAGACACCGC GTCTGTGGCG					1140
CTTAATGGCG	TGGGGTTAGA	TCCTTTAATA	TGTTATATAT	TCTGTTTCAT	CAATCACGTG	1200
GAATTACCGC	ACCCCAATCT	AGGAAATTAT	ACAATATATA	AGACAAAGTA	GTTAGTGCAC	
	TTTTGCAACC AAAACGTTGG					1260
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTCT	GTACCCTGAT	TGAGAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
	ATCCTGGTCA TAGGACCAGT					1380
GCTGCGCTTA	TAGTCTTGTG	TTTGTATGCC	TTTGTCCATT	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
	ATAAAGGTAG TATTTCCATC					1500
	GAAGCATTTA					1560
GGTTGTGGTC	CTTCGTAAAT	ACTCCTTTGC	GGTGTGTCGT	ACTGAATAAA	AGTTCTAACC	
	AATAAATAGT TTATTTATCA					1620
CACACTGGAA	TCAGTAGCCC	TTGAGCCATT	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTGA	1680
GTGTGACCTT	AGTCATCGGG	AACTCGGTAA	TTGTCGTCAC	AAGAAGACCG	TTCAAAAACT	
	AATGTATTCA TTACATAAGT					1740
ATCTCTATAG	CTCTGCTTCC	ТТСТАААТСА	AACCCATTGT	TGGATGCTCC	CTCTCCATTC	1800
TAGAGATATC	GAGACGAAGG	AAGATTTAGT	TTGGGTAACA	ACCTACGAGG	GAGAGGTAAG	

	TTGGCTTGCT AACCGAACGA			1860
	GTGTTATTTA CACAATAAAT			1920
	GTGCACATTT CACGTGTAAA			1980
	TGTGTTTATG ACACAAATAC			2040
	ACTAGATTAG TGATCTAATC			2100
	TAATGCTCCA ATTACGAGGT			2160
CGACAACAAC GCTGTTGTTG				

MVCGSPGGML	LLRAGLLALA	ALCLLRVPGA	RAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAIE	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PENLACEELP	VYDRGVCISP	EAIVTADGAD	FPMDSSNGNC	RGASSERCKC	180
KPIRATQKTY	FRNNYNYVIR	AKVKEIKTKC	HDVTAVVEVK	EILKSSLVNI	PRDTVNLYTS	240
SGCLCPPLNV	NEEYIIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLSK	300
SDSSNSDSTQ	SQKSGRNSNP	RQARN.				

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10. This sequence was assembled from public ESTs from the Genbank database (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

	GCCTTTTGGC					60
CCGCCTCGCC	CGGAAAACCG	CAGGTGACGC	GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
	GCAGCCCGGG					120
TACCAGACGC	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
	TGCTCCGGGT					180
CGAGAGACGG	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCMCMCCA	NO TO COMPAGE	CTCC1 1 C1 TC				
	AGTCCCTGCC					240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTCG	
ACTC ACCCCA	ACGCCATCCT	CCCCAMOCAC	03.00000033.00	omomoomooo	01.0001.0mgc	200
						300
IGAGICCGGI	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GTGGGTGACG	
AGCCCCGATC	TGCTCTTCTT	CCTCTCTCCC	AMCMA CCCCC	CC3 MCMCC3 C	CAMMO A CMMO	360
	ACGAGAAGAA					360
ICGGGGCIAG	ACGAGAAGAA	GGAGACACGG	TACATGCGCG	GGTAGACGTG	GTAACTGAAG	
CAGCACGAGC	CCATCAAGCC	ርጥርጥል እርጥርጥ	CTCTCCCACC	GGGCCCGGCA	ССССТСТСАС	420
	GGTAGTTCGG					420
	0011.011000	ORCHITCAGA	CACACGCICG	cccooccor	CCCGACACIC	
CCCATACTCA	TCAAGTACCG	CCACTCGTGG	CCGGAGAACC	TGGCCTGCGA	GGAGCTGCCA	480
	AGTTCATGGC					
					***************************************	
GTGTACGACA	GGGGCGTGTG	CATCTCTCCC	GAGGCCATCG	TTACTGCGGA	CGGAGCTGAT	540
	CCCCGCACAC					
TTTCCTATGG	ATTCTAGTAA	CGGAAACTGT	AGAGGGGCAA	GCAGTGAACG	CTGTAAATGT	600
AAAGGATACC	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTTACA	
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACTA	TGTCATTCGG	660
TTCGGATAAT	CTCGATGTGT	CTTCTGGATA	AAGGCCTTGT	TAATGTTGAT	ACAGTAAGCC	
GCTAAAGTTA	AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CTGCAGTAGT	GGAGGTGAAG	720
CGATTTCAAT	TTCTCTATTT	CTGATTCACG	GTACTACACT	GACGTCATCA	CCTCCACTTC	
	AGTCCTCTCT					780
CTCTAAGATT	TCAGGAGAGA	CCATTTGTAA	GGTGCCCTGT	GACAGTTGGA	GATATGGTCG	•
mamaaamaa	<b></b>					0.4.0
	TCTGCCCTCC					840
AGACCGACGG	AGACGGGAGG	TGAATTACAA	TTACTCCTTA	TATAGTAGTA	CCCGATACTT	

GATGAGGAAC GTTCCAGATT ACTCTTGGTG GAAGGCTCTA TAGCTGAGAA GTGGAAGGAT	
CTACTCCTTG CAAGGTCTAA TGAGAACCAC CTTCCGAGAT ATCGACTCTT CACCTTCCTA	900
CGACTCGGTA AAAAAGTTAA GCGCTGGGAT ATGAAGCTTC GTCATCTTGG ACTCAGTAAA	
GCTGAGCCAT TTTTTCAATT CGCGACCCTA TACTTCGAAG CAGTAGAACC TGAGTCATTT	960
AGTGATTCTA GCAATAGTGA TTCCACTCAG AGTCAGAAGT CTGGCAGGAA CTCGAACCCC	1020
TCACTAAGAT CGTTATCACT AAGGTGAGTC TCAGTCTTCA GACCGTCCTT GAGCTTGGGG	1020
CGGCAAGCAC GCAACTAAAT CCCGAAATAC AAAAAGTAAC ACAGTGGACT TCCTATTAAG GCCGTTCGTG CGTTGATTTA GGGCTTTATG TTTTTCATTG TGTCACCTGA AGGATAATTC	1080
ACTTACTTGC ATTGCTGGAC TAGCAAAGGA AAATTGCACT ATTGCACATC ATATTCTATT	
TGAATGAACG TAACGACCTG ATCGTTTCCT TTTAACGTGA TAACGTGTAG TATAAGATAA	1140
GTTTACTATA AAAATCATGT GATAACTGAT TATTACTTCT GTTTCTCTTT TGGTTTCTGC	1200
CAAATGATAT TTTTAGTACA CTATTGACTA ATAATGAAGA CAAAGAGAAA ACCAAAGACG	
TTCTCTCTTC TCTCAACCCC TTTGTAATGG TTTGGGGGGCA GACTCTTAAG TATATTGTGA	
AAGAGAGAAG AGAGTTGGGG AAACATTACC AAACCCCCGT CTGAGAATTC ATATAACACT	1260
GTTTTCTATT TCACTAATCA TGAGAAAAAC TGTTCTTTTG CAATAATAAT AAATTAAACA	1320
CAAAAGATAA AGTGATTAGT ACTCTTTTTG ACAAGAAAAC GTTATTATTA TTTAATTTGT	1320
TGCTGTTACC AGAGCCTCTT TGCTGAGTCT CCAGATGTTA ATTTACTTTC TGCACCCCAA ACGACAATGG TCTCGGAGAA ACGACTCAGA GGTCTACAAT TAAATGAAAG ACGTGGGGTT	1380
TTGGGAATGC AATATTGGAT GAAAAGAGA GTTTCTGGTA TTCACAGAAA GCTAGATATG	1440
AACCCTTACG TTATAACCTA CTTTTCTCTC CAAAGACCAT AAGTGTCTTT CGATCTATAC	1440
CCTTAAAACA TACTCTGCCG ATCTAATTAC AGCCTTATTT TTGTATGCCT TTTGGGCATT	1500
GGAATTTTGT ATGAGACGGC TAGATTAATG TCGGAATAAA AACATACGGA AAACCCGTAA	
CTCCTCATGC TTAGAAAGTT CCAAATGTTT ATAAAGGTAA AATGGCAGTT TGAAGTCAAA	
GAGGAGTACG AATCTTTCAA GGTTTACAAA TATTTCCATT TTACCGTCAA ACTTCAGTTT	1560
TGTCACATAG GCAAAGCAAT CAAGCACCAG GAAGTGTTTA TGAGGAAACA ACACCCAAGA	1620
ACAGTGTATC CGTTTCGTTA GTTCGTGGTC CTTCACAAAT ACTCCTTTGT TGTGGGTTCT	1020
TGAATTATTT TTGAGACTGT CAGGAAGTAA AATAAATAGG AGCTTAAGAA AGAACATTTT	
ACTTAATAAA AACTCTGACA GTCCTTCATT TTATTTATCC TCGAATTCTT TCTTGTAAAA	1680
GCCTGATTGA GAAGCACAAC TGAAACCAGT AGCCGCTGGG GTGTTAATGG TAGCATTCTT	1740
CGGACTAACT CTTCGTGTTG ACTTTGGTCA TCGGCGACCC CACAATTACC ATCGTAAGAA	1740
CTTTTGGCAA TACATTTGAT TTGTTCATGA ATATATTAAT CAGCATTAGA GAAATGAATT	1800
GAAAACCGTT ATGTAAACTA AACAAGTACT TATATAATTA GTCGTAATCT CTTTACTTAA	
ATAACTAGAC ATCTGCTGTT ATCACCATAG TTTTGTTTAA TTTGCTTCCT TTTAAATAAA	
TATTGATCTG TAGACGACAA TAGTGGTATC AAAACAAATT AAACGAAGGA AAATTTATTT	1860
CCCATTGGTG AAAGTCAAAA AAAAAAAAAA AAA	
GGGTAACCAC TTTCAGTTTT TTTTTTTTTT TTT	